

Table 2. *Halobacterium* NRC-1 genes and genetic elements. The classification heading is bolded and gene names (first column) and predicted gene products (second column) are listed.

Amino Acid Metabolism		Energy Metabolism	
<i>achY</i>	adenosylhomocysteinase	<i>speB</i>	agmatinase
<i>ansA</i>	L-asparaginase	<i>thrB</i>	homoserine kinase
<i>arcA</i>	arginine deiminase	<i>thrC1</i>	threonine synthase
<i>arcB</i>	ornithine carbamoyltransferase	<i>thrC2</i>	threonine synthase
<i>arcC</i>	carbamate kinase	<i>thrC3</i>	threonine synthase
<i>argB</i>	acetylglutamate kinase	<i>tmaA</i>	tryptophanase
<i>argG</i>	argininosuccinate synthetase	<i>trpA</i>	tryptophan synthase alpha chain
<i>argH</i>	argininosuccinate lyase	<i>trpB</i>	tryptophan synthase beta chain
<i>aroC</i>	chorismate synthase	<i>trpC</i>	indole-3-glycerol-phosphate synthase
<i>aroD</i>	3-dehydroquinate dehydratase	<i>trpD1</i>	phosphoribosyl transferase
<i>aroE</i>	shikimate 5-dehydrogenase	<i>trpD2</i>	phosphoribosyl transferase
<i>asd</i>	aspartate-semialdehyde dehydrogenase	<i>trpE1</i>	anthranilate synthase alpha chain
<i>asnA</i>	asparagine synthetase	<i>trpE2</i>	anthranilate synthase alpha chain
<i>aspB1</i>	aspartate aminotransferase	<i>trpF</i>	phosphoribosylanthranilate isomerase
<i>aspB2</i>	aspartate aminotransferase	<i>trpG1</i>	anthranilate synthase beta chain
<i>aspC1</i>	aspartate aminotransferase	<i>trpG2</i>	anthranilate synthase beta chain
<i>aspC2</i>	aspartate aminotransferase	<i>ykfB1</i>	chloromuconate cycloisomerase
<i>carA</i>	carbamoyl-phosphate synthase small subunit	<i>ykfB2</i>	chloromuconate cycloisomerase
<i>carB</i>	carbamoyl-phosphate synthase large subunit	<i>yrhA</i>	cysteine synthase
<i>cgs</i>	cystathione gamma-synthase	<i>yusM</i>	proline dehydrogenase
<i>cysK</i>	cysteine synthase		
<i>dapA</i>	dihydrodipicolinate synthase		
<i>gabD</i>	succinate-semialdehyde dehydrogenase		
<i>gabT</i>	gamma-aminobutyrate aminotransferase		
<i>gadD</i>	glutamate decarboxylase		
<i>gcvP1</i>	glycine dehydrogenase subunit 1		
<i>gcvP2</i>	glycine dehydrogenase subunit 2		
<i>gcvT1</i>	aminomethyltransferase		
<i>gcvT2</i>	aminomethyltransferase		
<i>gdhA1</i>	glutamate dehydrogenase		
<i>gdhA2</i>	glutamate dehydrogenase		
<i>gdhB</i>	glutamate dehydrogenase		
<i>glmS</i>	glutamine-fructose-6-phosphate transaminase		
<i>glnA</i>	glutamine synthetase		
<i>glyA</i>	glycine hydroxymethyltransferase		
<i>hakA</i>	kynureninase		
<i>hisA</i>	phosphoribosylformimino-5-aminoimidazole carboxamide ribotide isomerase		
<i>hisB</i>	imidazolylglycerol-phosphate dehydratase		
<i>hisC1</i>	histidinol-phosphate transaminase		
<i>hisC2</i>	histidinol-phosphate transaminase		
<i>hisD</i>	histidinol dehydrogenase		
<i>hisF</i>	imidazolylglycerol-phosphate synthase		
<i>hisG</i>	ATP phosphoribosyltransferase		
<i>hisH1</i>	imidazolylglycerol-phosphate synthase		
<i>hisH2</i>	imidazolylglycerol-phosphate synthase		
<i>hisJ</i>	phosphoribosyl-AMP cyclohydrolase		
<i>hisJ</i>	N1-(5-phospho-D-ribosyl)-AMP 1,6-hydrolase		
<i>hom</i>	homoserine dehydrogenase		
<i>hutG</i>	formiminoglutamate hydrolase		
<i>hutH</i>	histidine ammonia-lyase		
<i>hutI</i>	imidazolone-5-propionate hydrolase		
<i>hutU</i>	urocanate hydrolase		
<i>iluA</i>	threonine dehydratase		
<i>lysC</i>	aspartokinase II alpha subunit		
<i>mamA</i>	methylelaspartate mutase		
<i>mamB</i>	methylelaspartate mutase		
<i>metA</i>	homoserine O-acetyltransferase		
<i>metB</i>	cystathione alpha synthase		
<i>nadA</i>	quinolinate synthetase		
<i>nadB</i>	L-aspartate oxidase		
<i>ocd1</i>	ornithine cyclodeaminase		
<i>ocd2</i>	ornithine cyclodeaminase		
<i>pheA</i>	phenylalanine dehydratase		
<i>sat</i>	serine acetyltransferase		
<i>serA1</i>	phosphoglycerate dehydrogenase		
<i>serA2</i>	phosphoglycerate dehydrogenase		
<i>serA3</i>	phosphoglycerate dehydrogenase		
<i>serB</i>	phosphoserine phosphatase		
<i>soxB</i>	sarcosine oxidase		
Nucleotide Metabolism		Cofactor Metabolism	
<i>adk</i>	adenylate kinase	<i>birA</i>	biotin acetyl-CoA carboxylase ligase
<i>apt</i>	adenine phosphoribosyltransferase	<i>cbiA</i>	cobyric acid a,c-diamide synthase
<i>cda</i>	cytidine aminohydrolase	<i>cbiC</i>	precorrin isomerase
<i>cmk</i>	cytidylyl kinase	<i>cbiF</i>	cobalamin biosynthesis
<i>dcd</i>	deoxyctidyl triphosphate deaminase	<i>cbiG</i>	cobalamin biosynthesis
<i>dfp</i>	pantothenate metabolism flavoprotein	<i>cbiH</i>	cobalamin biosynthesis
<i>ddt</i>	deoxycytidine triphosphate deaminase	<i>cbiJ</i>	precorrin-3 methylase
<i>galE1</i>	UDP-glucose 4-epimerase	<i>cbiL</i>	cobalamin biosynthesis
<i>gptA1</i>	purine phosphoribosyltransferase	<i>cbiM</i>	cobalamin biosynthesis protein
<i>graD1</i>	glucose-1-phosphate thymidylyltransferase	<i>cbiP</i>	cobyric acid synthase
<i>graD2</i>	glucose-1-phosphate thymidylyltransferase	<i>cbiT</i>	cobalamin biosynthesis
<i>graD3</i>	glucose-1-phosphate thymidylyltransferase	<i>cobH</i>	cobalamin biosynthesis
<i>graD4</i>	glucose-1-phosphate thymidylyltransferase	<i>cobI</i>	cobalamin adenyltransferase
<i>graD5</i>	glucose-1-phosphate thymidylyltransferase	<i>cobN</i>	cobalamin biosynthesis protein
<i>graD6</i>	glucose-1-phosphate thymidylyltransferase	<i>ctaB</i>	heme synthase
<i>graD7</i>	glucose-1-phosphate thymidylyltransferase	<i>cysG</i>	uroporphyrin-III C-methyltransferase
<i>graD8</i>	glucose-1-phosphate thymidylyltransferase	<i>enB</i>	isochorismate
<i>graD9</i>	glucose-1-phosphate thymidylyltransferase	<i>gdb</i>	molybdopterin-guanine dinucleotide biosynthesis protein
<i>graD10</i>	glucose-1-phosphate thymidylyltransferase	<i>hem2</i>	porphobilinogen synthase
<i>graD11</i>	glucose-1-phosphate thymidylyltransferase	<i>hem3</i>	porphobilinogen deaminase
<i>graD12</i>	glucose-1-phosphate thymidylyltransferase	<i>hemK</i>	protoporphyrin oxidase
<i>graD13</i>	glucose-1-phosphate thymidylyltransferase	<i>hemL</i>	glutamate-1-semialdehyde aminotransferase
<i>graD14</i>	glucose-1-phosphate thymidylyltransferase	<i>hhoA</i>	4-hydroxybenzoate octaprenyltransferase
<i>graD15</i>	glucose-1-phosphate thymidylyltransferase	<i>hmcA</i>	protothymol IX magnesium chelatase
<i>graD16</i>	glucose-1-phosphate thymidylyltransferase	<i>hmoA</i>	molybdopterin oxidoreductase
<i>graD17</i>	glucose-1-phosphate thymidylyltransferase	<i>hxyA</i>	monoxygenase
<i>graD18</i>	glucose-1-phosphate thymidylyltransferase	<i>lipB</i>	lipoate protein ligase
<i>graD19</i>	glucose-1-phosphate thymidylyltransferase	<i>menA</i>	menaquinone biosynthesis
<i>graD20</i>	glucose-1-phosphate thymidylyltransferase	<i>menB</i>	dihydroxynaphthoic acid synthase
<i>graD21</i>	glucose-1-phosphate thymidylyltransferase	<i>menD</i>	2-succinyl-6-hydroxy-2,4-cyclohexadiene-1-carboxylate synthase
<i>graD22</i>	glucose-1-phosphate thymidylyltransferase	<i>menF</i>	isochorismate synthase
<i>graD23</i>	glucose-1-phosphate thymidylyltransferase	<i>moaA</i>	molybdenum cofactor biosynthesis protein
<i>graD24</i>	glucose-1-phosphate thymidylyltransferase	<i>moaB</i>	molybdenum cofactor biosynthesis protein
<i>graD25</i>	glucose-1-phosphate thymidylyltransferase	<i>moaC</i>	molybdenum cofactor biosynthesis
<i>graD26</i>	glucose-1-phosphate thymidylyltransferase	<i>moaE</i>	molybdenum cofactor biosynthesis
<i>graD27</i>	glucose-1-phosphate thymidylyltransferase	<i>moeA1</i>	molybdenum cofactor biosynthesis
<i>graD28</i>	glucose-1-phosphate thymidylyltransferase	<i>moeA2</i>	molybdenum cofactor biosynthesis
<i>graD29</i>	glucose-1-phosphate thymidylyltransferase	<i>moeB</i>	molybdenum cofactor biosynthesis
<i>graD30</i>	glucose-1-phosphate thymidylyltransferase	<i>nadC</i>	quinolinate phosphoribosyltransferase
<i>graD31</i>	glucose-1-phosphate thymidylyltransferase	<i>nadE</i>	NAD ⁺ synthetase
<i>graD32</i>	glucose-1-phosphate thymidylyltransferase	<i>nirD</i>	heme biosynthesis protein
<i>graD33</i>	glucose-1-phosphate thymidylyltransferase	<i>nirH</i>	heme biosynthesis protein
<i>graD34</i>	glucose-1-phosphate thymidylyltransferase	<i>nirJ</i>	heme biosynthesis protein
<i>graD35</i>	glucose-1-phosphate thymidylyltransferase	<i>pqqE</i>	coenzyme PQQ synthesis protein
<i>graD36</i>	glucose-1-phosphate thymidylyltransferase	<i>ribA</i>	GTP cyclohydrolase II
<i>graD37</i>	glucose-1-phosphate thymidylyltransferase	<i>ribC</i>	riboflavin synthase alpha subunit
		<i>ribE</i>	riboflavin synthase beta subunit
		<i>ribG</i>	riboflavin-specific deaminase
		<i>thi1</i>	thiamine biosynthetic enzyme
		<i>thiC</i>	thiamine biosynthesis protein
		<i>thiD</i>	hydroxymethylpyrimidine phosphate kinase
		<i>thiL</i>	thiamine monophosphate kinase
		<i>uroM</i>	S-adenosyl-L-methionine:uridylporphyrin III methyltransferase
		<i>yfkN</i>	2',3'-cyclic-nucleotide 2'-phosphodiesterase
		<i>yfmJ</i>	quinone oxidoreductase
Energy Metabolism			
		<i>aad</i>	aryl-alcohol dehydrogenase
		<i>acsJ</i>	acetyl-CoA synthetase
		<i>acs2</i>	acetyl-CoA synthetase
		<i>acs3</i>	acetyl-CoA synthetase
		<i>adh1</i>	alcohol dehydrogenase
		<i>adh2</i>	alcohol dehydrogenase
		<i>adh3</i>	alcohol dehydrogenase
		<i>adh4</i>	alcohol dehydrogenase
		<i>atpA</i>	H ⁺ -transporting ATP synthase subunit A
		<i>atpB</i>	H ⁺ -transporting ATP synthase subunit B
		<i>atpC</i>	H ⁺ -transporting ATP synthase subunit C
		<i>atpD</i>	H ⁺ -transporting ATP synthase subunit D
		<i>atpE</i>	H ⁺ -transporting ATP synthase subunit E
		<i>atpF</i>	H ⁺ -transporting ATP synthase subunit F
		<i>atpI</i>	H ⁺ -transporting ATP synthase subunit I
		<i>atpK</i>	H ⁺ -transporting ATP synthase subunit K
		<i>aup</i>	acetoin utilization protein
		<i>can</i>	aconitase
		<i>cat</i>	4-hydroxybutyrate CoA transferase
		<i>celM</i>	endoglucanase
		<i>citE</i>	citrate (pro-3S)-lyase
		<i>citZ</i>	citrate synthase
		<i>coxA1</i>	cytochrome c oxidase subunit I
		<i>coxA2</i>	cytochrome c oxidase subunit I
		<i>coxB1</i>	cytochrome c oxidase subunit II
		<i>coxB2</i>	cytochrome c oxidase subunit II
		<i>coxC</i>	cytochrome c oxidase subunit III
		<i>cyc</i>	cytochrome b ₆
		<i>cycD</i>	cytochrome d oxidase chain I
		<i>cydB</i>	cytochrome d oxidase chain II
		<i>deoC</i>	deoxyribose phosphate aldolase
		<i>dipS</i>	dihydrotereoate synthase
		<i>dld</i>	D-lactate dehydrogenase
		<i>dmsA</i>	dimethylsulfoxide reductase
		<i>dsa</i>	dihydrodipropioate S-acetyltransferase
		<i>eno</i>	phosphopyruvate hydratase
		<i>efA</i>	electron transfer flavoprotein subunit alpha
		<i>efB</i>	electron transfer flavoprotein subunit beta
		<i>fhp</i>	fructose-bisphosphatase
		<i>ferA1</i>	ferredoxin
		<i>ferA2</i>	ferredoxin
		<i>ferA3</i>	ferredoxin
		<i>ferB</i>	ferredoxin
		<i>fopL</i>	dihydrotereoate synthase
		<i>fprA</i>	flavoprotein
		<i>fucA</i>	fuculose-1-phosphate aldolase
		<i>fumC</i>	fumarate hydratase
		<i>gapA</i>	glyceraldehyde-3-phosphate dehydrogenase
		<i>gapB</i>	glyceraldehyde-3-phosphate dehydrogenase
		<i>gcd</i>	glucose dehydrogenase
		<i>gdcH</i>	glycine decarboxylase complex h-protein
		<i>glcC</i>	glycolate oxidase subunit
		<i>glcK</i>	glucose kinase
		<i>gldA</i>	sn-glycerol-1-phosphate dehydrogenase

<i>glpK</i>	glycerol kinase	<i>bop</i>	bacteriorhodopsin	<i>dppC1</i>	dipeptide ABC transporter permease	<i>ugpE</i>	sn-glycerol-3-phosphate transport system permease
<i>gpdA1</i>	glycerol-3-phosphate dehydrogenase chain A	<i>brp</i>	bacteriorhodopsin related protein	<i>dppC2</i>	dipeptide ABC transporter permease	<i>ybhF</i>	ABC-type transport protein
<i>gpdA2</i>	glycerol-3-phosphate dehydrogenase chain A	<i>crt</i>	carotenoid biosynthetic protein	<i>dppD</i>	dipeptide ABC transporter ATP-binding	<i>yfdM1</i>	ferrichrome ABC transporter permease protein
<i>gpdB</i>	glycerol-3-phosphate dehydrogenase chain B	<i>crtB1</i>	phytoene synthase	<i>dppF</i>	dipeptide ABC transporter ATP-binding	<i>yfdM2</i>	ferrichrome ABC transporter permease protein
<i>gpdC</i>	glycerol-3-phosphate dehydrogenase chain C	<i>crtB2</i>	phytoene synthase	<i>fepC</i>	ferric enterobactin transport protein	<i>yfmF</i>	ferrichrome ABC transporter ATP-binding protein
<i>gpm</i>	phosphoglycerate mutase	<i>crtI1</i>	phytoene dehydrogenase	<i>fhuG</i>	ferrichrome ABC transporter permease protein	<i>yfmO1</i>	multidrug resistance protein homolog
<i>hpcE</i>	2-hydroxyhepta-2,4-diene-1,7-dioate isomerase	<i>crtI2</i>	phytoene dehydrogenase	<i>ghpP</i>	proton/sodium-glutamate symport protein	<i>yfmO2</i>	multidrug resistance protein homolog
<i>icd</i>	isocitrate dehydrogenase, NADP	<i>crtI3</i>	phytoene dehydrogenase	<i>hemU</i>	iron (III) ABC transporter permease	<i>yhdG</i>	amino acid transporter
<i>ipp</i>	inorganic pyrophosphatase	<i>csg</i>	cell surface glycoprotein precursor	<i>hemV1</i>	iron (III) ABC transporter ATP-binding	<i>ykfD</i>	oligopeptide ABC transporter ATP-binding
<i>kdgK</i>	2-keto-3-deoxygluconate kinase	<i>dpg</i>	dolichol-P-glucosidase	<i>hemV2</i>	iron (III) ABC transporter ATP-binding	<i>ylnA</i>	anion permease
<i>korA</i>	putative 2-ketoglutarate ferredoxin oxidoreductase alpha	<i>est</i>	carboxylesterase	<i>ibp</i>	iron-binding protein	<i>yocR</i>	sodium-dependent transporter
<i>korB</i>	putative 2-ketoglutarate ferredoxin oxidoreductase beta	<i>exoM</i>	succinoglycan biosynthesis protein	<i>iucA</i>	iron transport protein A	<i>yagG</i>	phosphate ABC transporter binding
<i>lip</i>	lipoic acid synthase	<i>exsB</i>	succinoglycan biosynthesis	<i>iucB</i>	iron transport protein B	<i>yufN</i>	ABC transporter (lipoprotein)
<i>lpdA</i>	dihydrolipoamide dehydrogenase	<i>fabG</i>	3-oxoacyl-[acyl-carrier-protein] reductase	<i>iucC</i>	iron transport protein C	<i>yurY</i>	ABC transporter, ATP-binding protein
<i>mal</i>	methylaspartate ammonia-lyase	<i>fad1</i>	enoyl-CoA hydratase	<i>kdpA</i>	potassium-transferring ATPase A chain	<i>yusZ1</i>	oxidoreductase
<i>manC</i>	mannose-1-phosphate guanylyltransferase	<i>fad2</i>	enoyl-CoA hydratase	<i>kdpB</i>	potassium-transferring ATPase B chain	<i>yusZ2</i>	oxidoreductase
<i>mcmA1</i>	methylmalonyl-CoA mutase, subunit alpha	<i>fdfT</i>	farnesyldiphosphate farnesyltransferase	<i>kdpC</i>	potassium-transferring ATPase C chain	<i>yusZ3</i>	oxidoreductase
<i>mcmA2</i>	methylmalonyl-CoA mutase, subunit alpha	<i>fps</i>	putative isopentenyl pyrophosphate isomerase	<i>lctP</i>	L-lactate permease	<i>yvgX</i>	molybdenum-binding protein
<i>mcmA3</i>	methylmalonyl-CoA mutase, subunit alpha	<i>galE1</i>	UDP-D-mannose 4-epimerase	<i>nac</i>	sodium ⁺ - and chloride-dependent transporter	<i>yvrO</i>	amino acid ABC transporter, ATP-binding protein
<i>mdh</i>	malate dehydrogenase	<i>galE2</i>	UDP-glucose 4-epimerase	<i>nce</i>	$\text{Na}^+/\text{Ca}^{2+}$ -exchanging protein	<i>zntA</i>	zinc-transferring ATPase
<i>mdhA</i>	L-malate dehydrogenase	<i>gcdH</i>	glutaryl-CoA dehydrogenase	<i>nhaC1</i>	Na^+/H^+ antiporter	<i>zurA</i>	ABC transporter, ATP-binding protein
<i>mmdA</i>	methylmalonyl-CoA decarboxylase, subunit alpha	<i>ggt</i>	geranylgeranyl-diphosphate geranylgeranyltransferase	<i>nhaC2</i>	Na^+/H^+ antiporter	<i>zurM</i>	ABC transporter, permease protein
<i>moxR</i>	methanol dehydrogenase regulatory protein	<i>gmd</i>	GDP-D-mannose dehydratase	<i>nhaC3</i>	Na^+/H^+ antiporter		
<i>moz</i>	mybldopterin oxidoreductase	<i>hbd1</i>	3-hydroxyacyl-CoA dehydrogenase	<i>nifU</i>	nitrogen fixation protein		
<i>ndhG1</i>	NADH dehydrogenase/oxidoreductase	<i>hbd2</i>	3-hydroxyacyl-CoA dehydrogenase	<i>nosF1</i>	copper transport ATP-binding protein		
<i>ndhG2</i>	NADH dehydrogenase/oxidoreductase	<i>hop</i>	halorhodopsin	<i>nosF2</i>	copper transport ATP-binding protein		
<i>ndhG3</i>	NADH dehydrogenase/oxidoreductase	<i>idi</i>	isopentenyl pyrophosphate isomerase	<i>nosY</i>	nitrite/nitrate reduction protein		
<i>ndhG4</i>	NADH dehydrogenase/oxidoreductase	<i>ids</i>	bifunctional short chain isoprenyl diphosphate synthase	<i>oppD1</i>	oligopeptide ABC transporter ATP-binding		
<i>ndhG5</i>	NADH dehydrogenase/oxidoreductase	<i>idsA</i>	geranylgeranyl diphosphate synthase	<i>oppD2</i>	oligopeptide ABC transporter ATP-binding		
<i>nuoL</i>	F420H ₂ :quinone oxidoreductase chain L	<i>ilvE1</i>	branched-chain amino acid aminotransferase	<i>opuD</i>	glycine betaine transporter		
<i>nuoM</i>	F420H ₂ :quinone oxidoreductase chain M	<i>ilvE2</i>	branched-chain amino acid aminotransferase	<i>oxiT</i>	oxalate/formate antiporter		
<i>pdhA1</i>	pyruvate dehydrogenase alpha subunit	<i>lpb</i>	LPS biosynthesis protein	<i>panF</i>	pantothenate permease		
<i>pdhA2</i>	pyruvate dehydrogenase alpha subunit	<i>lpg</i>	LPS glycosyltransferase	<i>phmC</i>	phosphonates transport ATP-binding		
<i>pdhB</i>	pyruvate dehydrogenase beta subunit	<i>mvaA</i>	3-hydroxy-3-methylglutaryl-coenzyme A reductase	<i>phot1</i>	sodium dependent phosphate transporter		
<i>pgi</i>	glucose-6-phosphate isomerase	<i>mvaB</i>	3-hydroxy-3-methylglutaryl-coenzyme A synthase	<i>phot2</i>	sodium dependent phosphate transporter		
<i>pgk</i>	3-phosphoglycerate kinase	<i>mvk</i>	mevalonate kinase	<i>phot3</i>	sodium dependent phosphate transporter		
<i>pho2</i>	p-nitrophenyl phosphatase	<i>pan1</i>	membrane protein	<i>phoX</i>	phosphate ABC transporter periplasmic phosphatase-binding		
<i>pmm</i>	phosphoglucomutase/phospho-mannomutase	<i>pgsA</i>	CDP-diacylglycerol-glycerol-3-phosphate 3-phosphatidyltransferase	<i>potA1</i>	spermidine/putrescine ABC transporter ATP-binding		
<i>porA</i>	pyruvate ferredoxin oxidoreductase, subunit alpha	<i>pssA</i>	CDP-diacylglycerol-serine O-phosphatidyltransferase	<i>potA2</i>	spermidine/putrescine ABC transporter ATP-binding		
<i>porB</i>	pyruvate ferredoxin oxidoreductase, subunit beta	<i>rfbQ</i>	rhamsnosyl transferase	<i>potB</i>	spermidine/putrescine ABC transporter permease		
<i>ppiA</i>	peptidyl-prolyl isomerase	<i>sgb</i>	succinoglycan biosynthesis transport protein	<i>potC</i>	spermidine/putrescine ABC transporter permease		
<i>ppsa</i>	phosphoenolpyruvate synthase	<i>sop1</i>	sensory rhodopsin I	<i>potD</i>	spermidine/putrescine-binding protein		
<i>pykA</i>	pyruvate kinase	<i>sop2</i>	sensory rhodopsin II	<i>proX</i>	putative ABC transporter		
<i>rpi</i>	ribose 5-phosphate isomerase	<i>ugd</i>	UDP-glucose dehydrogenase	<i>psta1</i>	phosphate ABC transporter permease		
<i>sdhA</i>	succinate dehydrogenase subunit A	<i>ybjG</i>	membrane protein	<i>psta2</i>	phosphate ABC transporter ATP-binding		
<i>sdhB</i>	succinate dehydrogenase subunit B	<i>ycdH</i>	adhesion protein	<i>pstB1</i>	phosphate ABC transporter ATP-binding		
<i>sdhC</i>	succinate dehydrogenase subunit C			<i>pstB2</i>	phosphate ABC transporter ATP-binding		
<i>sucC</i>	succinyl-CoA synthetase beta chain			<i>pstC1</i>	phosphate ABC transporter permease		
<i>sucD</i>	succinyl-CoA synthetase alpha chain			<i>pstC2</i>	phosphate ABC transporter permease		
<i>suk</i>	sugar kinase			<i>puP</i>	proline permease		
<i>tpiA</i>	triosephosphate isomerase			<i>rbsA</i>	ribose ABC transporter ATP-binding		
<i>yafB</i>	aldehyde reductase			<i>rbsC1</i>	ribose ABC transporter permease		
<i>yjID</i>	NADH dehydrogenase			<i>rbsC2</i>	ribose ABC transporter permease		
<i>ykhA</i>	acyl-CoA hydrolase			<i>sfuB</i>	iron transporter-like protein		
<i>yngE</i>	propionyl-CoA carboxylase homolog			<i>trkA1</i>	TRK potassium uptake system protein		
<i>yqeC</i>	6-phosphogluconate dehydrogenase			<i>trkA2</i>	TRK potassium uptake system protein		
<i>yqjM</i>	NADH-dependent flavin oxidoreductase			<i>trkA3</i>	TRK potassium uptake system protein		
<i>ywfD</i>	glucose 1-dehydrogenase			<i>trkA4</i>	TRK potassium uptake system protein		
	Cell Envelope Components			<i>trkA5</i>	TRK potassium uptake system protein		
<i>acaB1</i>	3-ketoacyl-CoA thiolase	Transport		<i>trkA6</i>	TRK potassium uptake system protein		
<i>acaB2</i>	3-ketoacyl-CoA thiolase	<i>appB</i>	oligopeptide ABC permease	<i>trkH1</i>	TRK potassium uptake system protein		
<i>acc</i>	biotin carboxylase	<i>appC</i>	oligopeptide transport permease protein	<i>trkH2</i>	TRK potassium uptake system protein		
<i>acd1</i>	acyl-CoA dehydrogenase	<i>appF</i>	oligopeptide ABC transporter ATP-binding	<i>trkH3</i>	TRK potassium uptake system protein		
<i>acd2</i>	acyl-CoA dehydrogenase	<i>arsA1</i>	arsenical pump-driving ATPase	<i>trp1</i>	ABC transport protein		
<i>acd3</i>	acyl-CoA dehydrogenase	<i>arsA2</i>	arsenical pump-driving ATPase	<i>trp2</i>	ABC transport protein		
<i>acd4</i>	acyl-CoA dehydrogenase	<i>arsB</i>	arsenate transport protein	<i>trp3</i>	daunorubicin resistance ABC		
<i>acd5</i>	acyl-CoA dehydrogenase	<i>caa</i>	cation antiporter	<i>trp4</i>	transporter ATP-binding protein		
<i>act</i>	acyl-CoA thioester hydrolase	<i>cat1</i>	cationic amino acid transporter	<i>trp5</i>	ABC transporter, ATP-binding protein		
<i>aldY1</i>	aldehyde dehydrogenase (retinol)	<i>cat2</i>	cationic amino acid transporter	<i>trp6</i>	daunorubicin resistance ABC		
<i>aldY2</i>	aldehyde dehydrogenase (retinol)	<i>cat3</i>	cationic amino acid transporter	<i>trp7</i>	transporter ATP-binding protein		
<i>alkK</i>	medium-chain acyl-CoA ligase	<i>cat4</i>	cationic amino acid transporter	<i>trsE</i>	transfer complex protein		
<i>bchP</i>	geranylgeranyl hydrogenase	<i>cbn1</i>	cobalt transport protein	<i>ugpA</i>	sn-glycerol-3-phosphate transport system permease		
		<i>cbn2</i>	cobalt transport ATP-binding protein	<i>ugpB</i>	glycerol-3-phosphate-binding protein precursor		
		<i>cbn3</i>	cobalt transport protein	<i>ugpC</i>	sn-glycerol-3-phosphate transport system ATP-binding		

<i>gvpI1</i>	GvpI protein, cluster A	<i>hel</i>	DNA helicase	<i>rpoH</i>	DNA-directed RNA polymerase subunit H	<i>gatB1</i>	Glu-tRNA amidotransferase
<i>gvpI2</i>	GvpI protein, cluster B	<i>hjr</i>	holliday junction resolvase	<i>rpoK</i>	DNA-directed RNA polymerase subunit K	<i>gatB2</i>	Glu-tRNA amidotransferase
<i>gvpJ1</i>	GvpJ protein, cluster A	<i>hypA</i>	archaeal histone A1	<i>rpoL</i>	DNA-directed RNA polymerase subunit L	<i>gatC</i>	Glu-tRNA amidotransferase subunit C
<i>gvpJ2</i>	GvpJ protein, cluster B	<i>lig</i>	DNA ligase	<i>rpoM</i>	DNA-directed RNA-polymerase subunit M	<i>gltS</i>	glutamyl-tRNA synthetase
<i>gvpK1</i>	GvpK protein, cluster A	<i>mutL</i>	DNA mismatch repair protein	<i>rpoN</i>	DNA-directed RNA polymerase subunit N	<i>glyS</i>	glycine-tRNA synthetase
<i>gvpK2</i>	GvpK protein, cluster B	<i>mutS1</i>	mismatch repair protein	<i>snp</i>	snRNP homolog	<i>helA</i>	ATP-dependent helicase
<i>gvpL1</i>	GvpL protein, cluster A	<i>mutS2</i>	mismatch repair protein	<i>tbaP</i>	transcription initiation factor IID	<i>hemA</i>	glutamyl-tRNA reductase
<i>gvpL2</i>	GvpL protein, cluster B	<i>mutS3</i>	mismatch repair protein	<i>tbpB</i>	transcription initiation factor IID	<i>hepA</i>	ATP-dependent RNA helicase
<i>gvpM1</i>	GvpM protein, cluster A	<i>mutT</i>	mismatch repair protein	<i>tbpC</i>	transcription initiation factor IID	<i>hisS</i>	histidyl-tRNA synthetase
<i>gvpN1</i>	GvpN protein, cluster A	<i>mutY</i>	A/G specific adenine glycosylase, repair protein	<i>tbpD</i>	transcription initiation factor IID	<i>hypE</i>	hydrogenase expression/formation protein
<i>gvpN2</i>	GvpN protein, cluster B	<i>nfi</i>	endonuclease V	<i>tbpE</i>	transcription initiation factor IID	<i>ileS</i>	isoleucyl-tRNA synthetase
<i>gvpO1</i>	GvpO protein, cluster A	<i>nthA1</i>	endonuclease III	<i>tbpF</i>	transcription initiation factor IID	<i>infB</i>	bacterial-like IF2
<i>gvpO2</i>	GvpO protein, cluster B	<i>nthA2</i>	endonuclease III	<i>tbaA</i>	transcription initiation factor IIB	<i>ksgA</i>	dimethyladenosine transferase
<i>hacA</i>	atrazine chlorohydrolase	<i>nthB</i>	endonuclease III	<i>tbaB</i>	transcription initiation factor IIB	<i>leuS</i>	leucine-tRNA synthetase
<i>hsp1</i>	heat shock protein	<i>ogg</i>	8-oxoguanine DNA glycosylase	<i>tbaC</i>	transcription initiation factor IIB	<i>lysS</i>	lysyl-tRNA synthetase
<i>hsp2</i>	heat shock protein	<i>pcn</i>	proliferating-cell nuclear antigen	<i>tbaD</i>	transcription initiation factor IIB	<i>map</i>	methionyl aminopeptidase
<i>hsp3</i>	heat shock protein	<i>phr1</i>	photolyase/cryptochrome	<i>tbaE</i>	transcription initiation factor IIB	<i>metS</i>	methionine-tRNA synthetase
<i>hsp4</i>	heat shock protein	<i>phr2</i>	photolyase/cryptochrome	<i>tbaF</i>	transcription initiation factor IIB	<i>nop58</i>	archaeal nucleolar protein homolog
<i>hsp5</i>	heat shock protein	<i>pola1</i>	DNA polymerase type II small chain	<i>tbaG</i>	transcription initiation factor IIB	<i>pheS</i>	phenylalananyl-tRNA synthetase subunit alpha
<i>htr1</i>	Htr1 transducer	<i>pola2</i>	DNA polymerase type II large chain	<i>tbaH</i>	transcription initiation factor IIB	<i>pheY</i>	phenylalananyl-tRNA synthetase subunit beta
<i>htr2</i>	Htr2 transducer	<i>pob1</i>	DNA polymerase B1	<i>vacB</i>	ribonuclease II family protein	<i>pimT1</i>	L-isosoapartyl protein carboxyl methyltransferase
<i>htr3</i>	Htr3 transducer	<i>pob2</i>	DNA polymerase B2			<i>pimT2</i>	L-isosoapartyl-protein carboxyl methyltransferase
<i>htr4</i>	Htr4 transducer	<i>poc</i>	DNA polymerase bacteriophage-type			<i>proS</i>	proline-tRNA synthetase
<i>htr5</i>	Htr5 transducer	<i>pri</i>	DNA primase			<i>rfbU1</i>	LPS biosynthesis
<i>htr6</i>	Htr6 transducer	<i>rad2</i>	DNA repair protein			<i>rfbU2</i>	LPS biosynthesis
<i>htr7</i>	Htr7 transducer	<i>rad24a</i>	DNA repair protein			<i>rimI</i>	Pab N-terminal acetyltransferase
<i>htr8</i>	Htr8 transducer	<i>rad24b</i>	DNA repair protein			<i>rimK</i>	ribosomal protein S6 modification protein
<i>htr9</i>	Htr9 transducer	<i>rad24c</i>	DNA repair protein			<i>rnpB</i>	RNase P
<i>htr10</i>	Htr10 transducer	<i>rad25</i>	DNA repair protein			<i>rpl1p</i>	50S ribosomal protein L1P
<i>htr12</i>	Htr12 transducer	<i>rad3a</i>	helicase			<i>rpl2p</i>	50S ribosomal protein L2P
<i>htr13</i>	Htr13 transducer	<i>rad3b</i>	helicase			<i>rpl3p</i>	50S ribosomal protein L3P
<i>htr14</i>	Htr14 transducer	<i>radA1</i>	DNA repair protein			<i>rpl4e</i>	50S ribosomal protein L4E
<i>htr15</i>	Htr15 transducer	<i>radA2</i>	DNA repair protein			<i>rpl5p</i>	50S ribosomal protein L5P
<i>htr16</i>	Htr16 transducer	<i>repH</i>	replication protein			<i>rpl6p</i>	50S ribosomal protein L6P
<i>htr17</i>	Htr17 transducer	<i>repI</i>	replication protein			<i>rpl10e</i>	50S ribosomal protein L10E
<i>htr18</i>	Htr18 transducer	<i>repJ</i>	replication protein			<i>rpl10p</i>	50S ribosomal protein L10P
<i>htrA</i>	serine proteinase	<i>rfcA</i>	replication factor C small subunit			<i>rpl11p</i>	50S ribosomal protein L11P
<i>hyrA</i>	alkyl hydroperoxide reductase	<i>rfcC</i>	replication factor C small subunit			<i>rpl12p</i>	50S ribosomal protein L12P
<i>kaiC</i>	circadian regulator	<i>rfcB</i>	replication factor C large subunit			<i>rpl13p</i>	50S ribosomal protein L13P
<i>mcm</i>	MCM / cell division control protein 21	<i>rhl</i>	repair helicase			<i>rpl14p</i>	50S ribosomal protein L14P
<i>merA</i>	mercury(II) reductase	<i>rmeM</i>	type I restriction-modification enzyme, M subunit			<i>rpl15e</i>	50S ribosomal protein L15E
<i>minD1</i>	cell division inhibitor	<i>rmeS</i>	type I restriction-modification enzyme, S subunit			<i>rpl15p</i>	50S ribosomal protein L15P
<i>minD2</i>	cell division inhibitor	<i>rmeR</i>	type I restriction-modification enzyme, R subunit			<i>rpl18e</i>	50S ribosomal protein L18E
<i>noxA</i>	NADH oxidase	<i>rnh</i>	RNAse H			<i>rpl18p</i>	50S ribosomal protein L18P
<i>noxC</i>	NADH oxidase	<i>rpa</i>	replication A related protein			<i>rpl19e</i>	50S ribosomal protein L19E
<i>orc1</i>	orc / cell division control protein 6	<i>sod1</i>	superoxide dismutase			<i>rpl21e</i>	50S ribosomal protein L21E
<i>orc2</i>	orc / cell division control protein 6	<i>sod2</i>	superoxide dismutase			<i>rpl22p</i>	50S ribosomal protein L22P
<i>orc3</i>	orc / cell division control protein 6	<i>ssrA</i>	integrase/recombinase			<i>rpl23p</i>	50S ribosomal protein L23P
<i>orc4</i>	orc / cell division control protein 6	<i>top6A</i>	DNA topoisomerase VI subunit A			<i>rpl24e</i>	50S ribosomal protein L24E
<i>orc5</i>	orc / cell division control protein 6	<i>top6B</i>	DNA topoisomerase VI subunit B			<i>rpl24p</i>	50S ribosomal protein L24P
<i>orc6</i>	orc / cell division control protein 6	<i>topA</i>	DNA topoisomerase I			<i>rpl29p</i>	50S ribosomal protein L29P
<i>orc7</i>	orc / cell division control protein 6	<i>uvrA</i>	excision nuclease chain A			<i>rpl30p</i>	50S ribosomal protein L30P
<i>orc8</i>	orc / cell division control protein 6	<i>uvrB</i>	excision nuclease chain B			<i>rpl31e</i>	50S ribosomal protein L31E
<i>orc9</i>	orc / cell division control protein 6	<i>uvrC</i>	excision nuclease chain C			<i>rpl32e</i>	50S ribosomal protein L32E
<i>pelA</i>	cell division protein pelota	<i>uvrD</i>	repair helicase			<i>rpl37e</i>	50S ribosomal protein L37E
<i>perA</i>	peroxidase / catalase	<i>xthA</i>	endonuclease IV			<i>rpl39e</i>	50S ribosomal protein L39E
<i>psmA</i>	proteasome, subunit alpha	<i>yhcR</i>	phosphoesterase			<i>rpl40e</i>	50S ribosomal protein L40E
<i>psmB</i>	proteasome, subunit beta	<i>yqjH</i>	trans lesion repair			<i>rpl44e</i>	50S ribosomal protein L44E
<i>rrt</i>	7s RNA	<i>zim</i>	CTAG modification methylase			<i>rps2p</i>	30S ribosomal protein S2P
<i>rspA</i>	starvation sensing protein					<i>rps3e</i>	30S ribosomal protein S3E
<i>sec11</i>	signal sequence peptidase					<i>rps3p</i>	30S ribosomal protein S3P
<i>secD</i>	protein-export membrane protein					<i>rps4e</i>	30S ribosomal protein S4E
<i>secE</i>	protein translocase					<i>rps4p</i>	30S ribosomal protein S4P
<i>secF</i>	protein-export membrane protein					<i>rps5p</i>	30S ribosomal protein S5P
<i>secY</i>	protein translocase					<i>rps6e</i>	30S ribosomal protein S6E
<i>smc1</i>	chromosome segregation	<i>brr2</i>	pre-mRNA splicing helicase			<i>rphs6</i>	30S ribosomal protein HS6
<i>sojA</i>	SpoOA activation inhibitor	<i>cca</i>	tRNA nucleotidyltransferase			<i>rps7p</i>	30S ribosomal protein S7P
<i>sojB</i>	SpoOA activation inhibitor	<i>endA</i>	tRNA intron endonuclease			<i>rps8e</i>	30S ribosomal protein S8E
<i>sojC1</i>	SpoOA activation inhibitor	<i>epf1</i>	mRNA 3'-end processing factor homolog			<i>rps8p</i>	30S ribosomal protein S8P
<i>sojC2</i>	SpoOA activation inhibitor	<i>epf2</i>	mRNA 3'-end processing factor homolog			<i>rps9p</i>	30S ribosomal protein S9P
<i>sojD</i>	SpoOA activation inhibitor	<i>nusA</i>	transcription termination-antitermination factor			<i>rps10p</i>	30S ribosomal protein S10P
<i>sojE</i>	SpoOA activation inhibitor	<i>nusG</i>	transcription termination-antitermination factor			<i>rps11p</i>	30S ribosomal protein S11P
<i>srp19</i>	signal recognition particle	<i>rpb3</i>	DNA-directed RNA polymerase II			<i>rps12p</i>	30S ribosomal protein S12P
<i>sdp54</i>	signal recognition particle	<i>rpoA</i>	DNA-directed RNA polymerase subunit A			<i>rps13p</i>	30S ribosomal protein S13P
<i>surE</i>	stationary phase survival protein	<i>rpoB'</i>	DNA-directed RNA polymerase subunit B'			<i>rps14p</i>	30S ribosomal protein S14P
<i>tatC1</i>	protein export	<i>rpoB''</i>	DNA-directed RNA polymerase subunit B''			<i>rps15p</i>	30S ribosomal protein S15P
<i>tatC2</i>	protein export	<i>rpoC</i>	DNA-directed RNA polymerase subunit C			<i>rps17e</i>	30S ribosomal protein S17E
<i>trzA</i>	N-ethylammeline chlorohydrolase	<i>rpoE'</i>	DNA-directed RNA polymerase subunit E'			<i>rps17p</i>	30S ribosomal protein S17P
		<i>rpoE''</i>	DNA-directed RNA polymerase subunit E''			<i>rps19e</i>	30S ribosomal protein S19E
						<i>rps19p</i>	30S ribosomal protein S19P
						<i>rps24e</i>	30S ribosomal protein S24E
						<i>rps27ae</i>	30S ribosomal protein S27AE
						<i>rps27e</i>	30S ribosomal protein S27E
DNA Replication, Repair, and Recombination							
<i>alkA</i>	3-methyladenine DNA glycosylase						
<i>dinF</i>	DNA damage-inducible protein						
<i>dip1</i>	DNA damage-inducible protein						
<i>dip2</i>	DNA damage-inducible protein						
<i>dip3</i>	competence-damage protein CinA-like						
<i>gyrA</i>	DNA gyrase subunit A						
<i>gyrB</i>	DNA gyrase subunit B						
Transcription							
<i>brr2</i>	pre-mRNA splicing helicase						
<i>cca</i>	tRNA nucleotidyltransferase						
<i>endA</i>	tRNA intron endonuclease						
<i>epf1</i>	mRNA 3'-end processing factor homolog						
<i>epf2</i>	mRNA 3'-end processing factor homolog						
<i>nusA</i>	transcription termination-antitermination factor						
<i>nusG</i>	transcription termination-antitermination factor						
<i>rpb3</i>	DNA-directed RNA polymerase II						
<i>rpoA</i>	DNA-directed RNA polymerase subunit A						
<i>rpoB'</i>	DNA-directed RNA polymerase subunit B'						
<i>rpoB''</i>	DNA-directed RNA polymerase subunit B''						
<i>rpoC</i>	DNA-directed RNA polymerase subunit C						
<i>rpoE'</i>	DNA-directed RNA polymerase subunit E'						
<i>rpoE''</i>	DNA-directed RNA polymerase subunit E''						
Translation							
<i>alaS</i>	alanyl-tRNA synthetase						
<i>argS</i>	arginine-tRNA synthetase						
<i>aspS</i>	aspartyl-tRNA synthetase						
<i>cra</i>	proliferating-cell nucleolar antigen						
<i>cysS</i>	cysteinyl-tRNA synthetase						
<i>drg</i>	GTP-binding protein DRG						
<i>dys</i>	deoxyhypusine synthase						
<i>eef1a</i>	translation elongation factor eEF-1A						
<i>eef1b</i>	translation elongation factor eEF-1A subunit alpha						
<i>eef1f</i>	translation elongation factor eEF-1 subunit beta						
<i>eef2</i>	translation elongation factor eEF-2						
<i>eif1a1</i>	translation initiation factor eIF-1A						
<i>eif1a2</i>	translation initiation factor eIF-1A						
<i>eif2a</i>	translation initiation factor eIF-2						
<i>eif2b</i>	translation initiation factor eIF-2 subunit alpha						
<i>eif2d</i>	translation initiation factor eIF-2 subunit beta						
<i>eif2ba</i>	translation initiation factor eIF-2B subunit alpha						
<i>eif2bd</i>	translation initiation factor eIF-2B subunit delta						
<i>eif2g</i>	translation initiation factor eIF-2 subunit gamma						
<i>eif4a</i>	ATP-dependent RNA helicase homolog eIF-4A						
<i>eif5a</i>	translation initiation factor eIF-5A						
<i>erf1</i>	peptide chain release factor eRF-1						
<i>fib</i>	fibrillarin						
<i>gatA</i>	Glu-tRNA amidotransferase						

<i>rps28e</i>	30S ribosomal protein S28E	<i>valS</i>	valyl-tRNA synthetase	<i>hit1</i>	histidine triad protein	<i>oxrA</i>	putative oxidoreductase
<i>rrlA</i>	23S rRNA			<i>hit2</i>	histidine triad protein	<i>panB</i>	ketopantoate hydroxymethyltransferase-like
<i>rrlB</i>	5s rRNA			<i>hip</i>	hemolysin protein	<i>pca</i>	plastocyanin homology
<i>rrs</i>	16S rRNA			<i>hly</i>	halolysin	<i>pchA</i>	potassium channel homolog
<i>serS</i>	serine-tRNA synthetase	<i>aca</i>	probable acetyl-coa acetyltransferase	<i>hmp</i>	putative membrane protein	<i>pchB</i>	potassium channel homolog
<i>sus</i>	tRNA-pseudouridine synthase	<i>apa</i>	diadenosine tetraphosphate	<i>hpb</i>	possible phosphate binding protein	<i>pepB1</i>	aminopeptidase homolog
<i>tgtA1</i>	archaeosine tRNA-ribosyltransferase	<i>aph</i>	pyrophosphohydrolase	<i>htlA</i>	Htr-like protein	<i>pepB2</i>	aminopeptidase homolog
<i>tgtA2</i>	archaeosine tRNA-ribosyltransferase	<i>appA</i>	oligopeptide binding protein	<i>htlB</i>	Htr-like protein	<i>pepB3</i>	aminopeptidase homolog
<i>thrS</i>	threonyl-tRNA synthetase	<i>araL</i>	unknown L-arabinose operon protein	<i>htlC</i>	Htr-like protein	<i>pepQ1</i>	probable peptidase
<i>trm1</i>	N2,N2-dimethylguanosine tRNA methyltransferase	<i>atp</i>	aaα superfamily atpase	<i>htlD</i>	Htr-like protein	<i>pepQ2</i>	probable peptidase
<i>trn1</i>	Phe-tRNA-GAA	<i>bcp</i>	bacterioferritin comigrating protein	<i>icfa</i>	carbonic anhydrase	<i>pgp</i>	phosphoglycolate phosphatase
<i>trn2</i>	Val-tRNA-TAC	<i>bdb</i>	L-2,4-diaminobutyrate decarboxylase	<i>imp</i>	immunogenic protein	<i>phnE</i>	transport protein
<i>trn3</i>	Arg-tRNA-GCG	<i>bih</i>	Brp-like homolog	<i>imb</i>	oxidoreductase homolog	<i>phnF</i>	
<i>trn4</i>	Asn-tRNA-GTT	<i>blp</i>	bacterio-opsin linked product	<i>ISH1</i>	insertion sequence	<i>pmu1</i>	phosphomannomutase
<i>trn5</i>	Ile-tRNA-CAT	<i>boa1</i>	bacterio-opsin activator-like protein	<i>ISH2</i>	insertion sequence (same IRs as ISH8)	<i>pmu2</i>	phosphomannomutase
<i>trn6</i>	Tyr-tRNA-GTA	<i>boa2</i>	bacterio-opsin activator-like protein	<i>ISH3</i>	insertion sequence family (similar to ISH27 and ISH51)	<i>pnm</i>	N-methyltransferase homolog
<i>trn7</i>	Ser-tRNA-CGA	<i>boa3</i>	bacterio-opsin activator-like protein	<i>ISH4</i>	insertion sequence	<i>ppd</i>	3-isopropylmalate dehydratase
<i>trn8</i>	Gln-tRNA-TTG	<i>boa4</i>	bacterio-opsin activator-like protein	<i>ISH5</i>	insertion sequence family	<i>ppe</i>	phosphoesterase
<i>trn9</i>	Met-tRNA-initiation	<i>caaX</i>	zinc metalloproteinase homolog	<i>ISH6</i>	insertion sequence (similar to ISH51)	<i>prkA</i>	protein kinase
<i>trn10</i>	Arg-tRNA-TCG	<i>cad</i>	pterin-4a-carbinolamine dehydratase	<i>ISH7</i>	insertion sequence family (similar to ISH24)	<i>prp2</i>	phosphate regulatory protein homolog
<i>trn11</i>	Ala-tRNA-GGC	<i>cbp</i>	calcium-binding protein homology	<i>ISH8</i>	insertion sequence family (same IRs as ISH2, similar to ISH26)	<i>prrIV1</i>	ATP-dependent 26S proteinase
<i>trn12</i>	Arg-tRNA-CCT	<i>ccp</i>	cytochrome aa3 controlling protein	<i>ISH9</i>	insertion sequence (similar to ISH28)	<i>prrIV2</i>	regulatory subunit 4 homolog
<i>trn13</i>	Gly-tRNA-TCC	<i>chi</i>	chitinase	<i>ISH10</i>	insertion sequence	<i>psc</i>	3-phosphoshikimate 1-carboxyvinyltransferase
<i>trn14</i>	Lys-tRNA-CTT	<i>comA</i>	competence-like protein	<i>ISH11</i>	insertion sequence	<i>ptp</i>	probable transport protein
<i>trn15</i>	Trp-tRNA-CCA	<i>cpeE</i>	phycocyanin alpha phycocyanobilin lyase-like	<i>ISH12</i>	insertion sequence family (similar to ISH1.8)	<i>rli</i>	RNAse L inhibitor homolog
<i>trn16</i>	Thr-tRNA-CGT	<i>cpx</i>	probable carboxypeptidase	<i>kinA1</i>	signal-transducing histidine kinase homolog	<i>scm</i>	24-sterol C-methyltransferase
<i>trn17</i>	Ser-tRNA-GCT	<i>cyo</i>	cytochrome oxidase subunit I homolog	<i>kinA2</i>	signal-transducing histidine kinase homolog	<i>sdh</i>	succinate dehydrogenase complex protein
<i>trn18</i>	Leu-tRNA-CAG	<i>dbp</i>	DNA binding protein eukaryotic-like	<i>lds</i>	long hypothetical diphthine synthase	<i>sdhD</i>	succinate dehydrogenase membrane
<i>trn19</i>	Thr-tRNA-TCT	<i>dgs</i>	dolichol-P-glucose transferase	<i>lfl1</i>	long-chain fatty-acid-CoA ligase	<i>slyD</i>	peptidyl-prolyl cis-trans isomerase
<i>trn20</i>	Ser-tRNA-TGA	<i>edp</i>	proteinas IV homolog	<i>lfl2</i>	long-chain fatty-acid-CoA ligase	<i>spoVR</i>	spore cortex synthesis protein
<i>trn21</i>	Glu-tRNA-TTC	<i>eye</i>	erythromycin esterase-like	<i>lon</i>	ATP-dependent proteinase homolog	<i>sub</i>	subtilisin homolog
<i>trn22</i>	Glu-tRNA-CTC	<i>fbr</i>	cytochrome-like protein	<i>lpl</i>	lipoate protein ligase	<i>suhB</i>	extragenic suppressor homolog
<i>trn23</i>	Leu-tRNA-CAA	<i>fold</i>	methylenetetrahydrofolate dehydrogenase	<i>lta</i>	l-allo-threonine aldolase	<i>tagD</i>	glycerol-3-phosphate cytidyltransferase
<i>trn24</i>	Leu-tRNA-TAG			<i>mak</i>	MAPK-activated protein kinase	<i>thb</i>	thiamin-binding periplasmic protein precursor homolog
<i>trn25</i>	Leu-tRNA-GAG	<i>gbp1</i>	GTP-binding protein homolog	<i>maoC1</i>	monoamine oxidase regulatory-like	<i>tmp</i>	putative transposase
<i>trn26</i>	Val-tRNA-CAC	<i>gbp2</i>	GTP-binding protein homolog	<i>maoC2</i>	monoamine oxidase regulatory-like	<i>tot</i>	transmembrane oligosaccharyl transferase homolog
<i>trn27</i>	Met-tRNA-CAT	<i>gbp3</i>	GTP-binding protein homolog	<i>mch</i>	N5,N10-methenyltetrahydromanopterin cyclohydrolase	<i>tpc</i>	RNA 3'-terminal phosphate cyclase
<i>trn28</i>	Val-tRNA-GAC	<i>gbp4</i>	GTP-binding protein homolog	<i>mer</i>	N5,N10-methylenetetrahydromanopterin reductase	<i>trab</i>	possible signaling protein
<i>trn29</i>	Gly-tRNA-GCC	<i>gbp5</i>	GTP-binding protein homolog	<i>mrr</i>	Mrr restriction system protein homolog	<i>trh</i>	thioredoxin reductase-like protein
<i>trn30</i>	Pro-tRNA-GGG	<i>gcp</i>	O-sialoglycoprotein endopeptidase homolog	<i>msrA</i>	peptide methionine sulfoxide reductase	<i>tssA</i>	probable thiosulfate sulfurtransferase
<i>trn31</i>	Ile-tRNA-GAT	<i>glo1</i>	glyoxalase	<i>nifS</i>	NifS protein, class-V aminotransferase	<i>tssB</i>	thiosulfate sulfurtransferase
<i>trn32</i>	Leu-tRNA-TAA	<i>gst</i>	galactosyltransferase homolog	<i>nodP</i>	nodulation protein	<i>uae</i>	UDP-N-acetylglucosamine 2-epimerase
<i>trn33</i>	Pro-tRNA-CGG	<i>glt</i>	glycosyl transferase-like	<i>nodA</i>	NADH dehydrogenase/oxidoreductase-like protein	<i>yajO1</i>	probable oxidoreductase
<i>trn34</i>	Gly-tRNA-CCC	<i>gufA</i>	GufA protein	<i>nodB</i>	NADH dehydrogenase/oxidoreductase-like protein	<i>yajO2</i>	probable oxidoreductase
<i>trn35</i>	Thr-tRNA-GGT	<i>gull</i>	inosine-5'-monophosphate dehydrogenase-like	<i>nodC</i>	NADH dehydrogenase/oxidoreductase-like protein	<i>yuiH</i>	sulfite oxidase homolog
<i>trn36</i>	Ser-tRNA-GGA	<i>gul2</i>	inosine-5'-monophosphate dehydrogenase-like	<i>nodD</i>	NADH dehydrogenase/oxidoreductase-like protein	<i>yuxL</i>	acylaminocycl-peptidase
<i>trn37</i>	His-tRNA-GTG	<i>hal</i>	O-acetyl homoserine	<i>npa</i>	neutral proteinase	<i>yvbT</i>	alkanal monooxygenase homolog
<i>trn38</i>	Pro-tRNA-TGG	<i>ham1</i>	HAM1 protein homolog	<i>ntp</i>	neutral proteinase	<i>yvoF</i>	acyltransferase homolog
<i>trn39</i>	Gln-tRNA-CTG	<i>hat1</i>	acetyltransferase homolog			<i>ywaD</i>	aminopeptidase
<i>trn40</i>	Arg-tRNA-TCT	<i>hat2</i>	acetyltransferase homolog			<i>yyaI</i>	acyltransferase homolog
<i>trn41</i>	Ala-tRNA-TGC	<i>hcpA</i>	halocyanin precursor-like				
<i>trn42</i>	Cys-tRNA-GCA	<i>hcpB</i>	halocyanin precursor-like				
<i>trn43</i>	Lys-tRNA-TTT	<i>hcpC</i>	halocyanin precursor-like				
<i>trn44</i>	Arg-tRNA-CCG	<i>hcpD</i>	halocyanin precursor-like				
<i>trn45</i>	Ala-tRNA-CGC	<i>hdrD</i>	heterodisulfide reductase				
<i>trn46</i>	Ala-tRNA-CGC	<i>hef</i>	hydrogenase expression/formation				
<i>trn47</i>	Asp-tRNA-GTC	<i>hflX1</i>	GTP-binding protein				
<i>trpS1</i>	tryptophanyl-tRNA synthetase	<i>hflX2</i>	GTP-binding protein				
<i>trpS2</i>	tryptophanyl-tRNA synthetase						
<i>truA</i>	pseudouridylate synthase I						
<i>tyrS</i>	tyrosyl-tRNA synthetase						